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<141> 2006-06-14
<150> PCT/EP04/014266
<151> 2004-12-15
<150> DE 10359594.5
<151> 2003-12-18
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Asn Ala Ile Glu Gly Ile Phe Gln Thr Asp Arg Gly Phe Thr His Thr 65 70 75 80

Val Ala Thr Gln Val Gly Ala Trp Glu Gln Gln Met Ala Met Lys Gly 85 90 95

Lys His Val Lys Pro Ala Phe Asp Asp Thr Leu Glu Gly Phe Glu Tyr 100 105 110

Val Leu Asp Ala Val Thr Gly Arg Thr Pro Ile Ser Gln Gln Trp Ile 115 120 125

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- Ala Leu Arg Pro Thr Arg Val Leu Pro Glu Leu His Asp Ala Ala His 305 310 315 320
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- Glu Gly Lys Arg Asn Gly Leu Glu Phe Leu Leu Gln Arg Ile Phe Ile 340 345 350
- Gly Ser Pro Phe Asn Leu Pro Glu Gly Tyr Asn Ala Phe Pro Asp Ser 355 360 365
- Tyr Cys Leu Thr Leu Ala Phe Asn Ser Asn Ser Pro Lys Gln Ile Phe 370 375 380
- His Pro Leu Ser Ile Val Ile Ala Ala Arg Asp Gly Lys Arg Ala Ser 385 390 395 400
- Ser Asp Leu Val Ala Ala Thr Ser Ile Gly Tyr Asn Phe His Ala Tyr 405 410 415
- Gly Arg Glu Val Glu Pro Val Val Thr Glu Ser Phe Arg Glu Arg Val 420 425 430
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<212> DNA
<213> Artificial Sequence
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